

SEQUENCE LISTING

<110> KROHN, KAI
HEINO, MAARIT
PETERSON, PART
SCOTT, HAMISH S
ANTONARAKIS, STYLIANOS E
LALIOTI, MARIA D
SHIMIZU, NOBUYOSHI D
KUDOH, JUN D

D) ><120> NOVEL GENE DEFECTIVE IN APECED AND ITS USE

<130> U 012653-9

<140> 09/508,658
<141> 2000-11-03

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<151> 1998-09-23

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<170> PatentIn version 3.1

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PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re application of: Kai Krohn, et al.

Serial No.: 09/508,658

Group No.: 1634

Filed: November 3, 2000

Examiner.: Diana B. Johannsen

For: NOVEL GENE DEFECTIVE IN APECED AND ITS USE

Attorney Docket No.: U 012653-9

Assistant Commissioner of Patents
Washington, D.C. 20231

AMENDMENT

Please replace the sequence listing on file in this application with the attached sequence listing. Support for SEQ ID NO: 3 to 7 is found in Figure 5 and on page 6, lines 4-9 of the specification.

Respectfully submitted,

Janet I. Cord
c/o Ladas & Parry
26 West 61st Street
New York, NY 10023
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CERTIFICATE OF MAILING /TRANSMISSION(37 CFR 1.8a)

I hereby certify that this correspondence is, on the date shown below, being:

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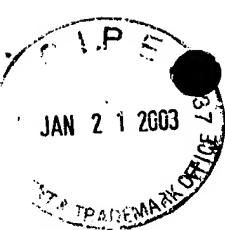
transmitted by facsimile to the Patent and Trademark Office to fax number (703) 308-7922.

Signature

JANET CORD

(type or print name of person certifying)

Date: January 15, 2003



SEQUENCE LISTING

<110> KROHN, KAI
HEINO, MAARIT
PETERSON, PART
SCOTT, HAMISH S
ANTONARAKIS, STYLIANOS E
LALIOTI, MARIA D
SHIMIZU, NOBUYOSHI D
KUDOH, JUN D

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Ile Leu Ile Gln Gln Val Phe Glu Ser Gly Gly Ser Lys Lys Cys Ile
210 215 220

Gln Val Gly Gly Glu Phe Tyr Thr Pro Ser Lys Phe Glu Asp Ser Gly
225 230 235 240

Ser Gly Lys Asn Lys Ala Arg Ser Ser Gly Pro Lys Pro Leu Val
245 250 255

Arg Ala Lys Gly Ala Gln Gly Ala Ala Pro Gly Gly Glu Ala Arg
260 265 270

Leu Gly Gln Gln Gly Ser Val Pro Ala Pro Leu Ala Leu Pro Ser Asp
275 280 285

Pro Gln Leu His Gln Lys Asn Glu Asp Glu Cys Ala Val Cys Arg Asp
290 295 300

Gly Gly Glu Leu Ile Cys Cys Asp Gly Cys Pro Arg Ala Phe His Leu
305 310 315 320

Ala Cys Leu Ser Pro Pro Leu Arg Glu Ile Pro Ser Gly Thr Trp Arg

325

330

335

Cys Ser Ser Cys Leu Gln Ala Thr Val Gln Glu Val Gln Pro Arg Ala
340 345 350

Glu Glu Pro Arg Pro Gln Glu Pro Pro Val Glu Thr Pro Leu Pro Pro
355 360 365

Gly Leu Arg Ser Ala Gly Glu Glu Val Arg Gly Pro Pro Gly Glu Pro
370 375 380

Leu Ala Gly Met Asp Thr Thr Leu Val Tyr Lys His Leu Pro Ala Pro
385 390 395 400

Pro Ser Ala Ala Pro Leu Pro Gly Leu Asp Ser Ser Ala Leu His Pro
405 410 415

Leu Leu Cys Val Gly Pro Glu Gly Gln Gln Asn Leu Ala Pro Gly Ala
420 425 430

Arg Cys Gly Val Cys Gly Asp Gly Thr Asp Val Leu Arg Cys Thr His
435 440 445

Cys Ala Ala Ala Phe His Trp Arg Cys His Phe Pro Ala Gly Thr Ser
450 455 460

Arg Pro Gly Thr Gly Leu Arg Cys Arg Ser Cys Ser Gly Asp Val Thr
465 470 475 480

Pro Ala Pro Val Glu Gly Val Leu Ala Pro Ser Pro Ala Arg Leu Ala
485 490 495

Pro Gly Pro Ala Lys Asp Asp Thr Ala Ser His Glu Pro Ala Leu His
500 505 510

Arg Asp Asp Leu Glu Ser Leu Leu Ser Glu His Thr Phe Asp Gly Ile
515 520 525

Leu Gln Trp Ala Ile Gln Ser Met Ala Arg Pro Ala Ala Pro Phe Pro
530 535 540

Ser
545

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Pro Arg Ala Phe His Leu Ala Cys Leu Ser Pro Pro Leu Arg Glu Ile
20 25 30

Pro Ser Gly Thr Trp Arg Cys Ser Ser Cys
35 40

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<212> PRT
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Ala Ala Ala Phe His Trp Arg Cys His Phe Pro Ala Gly Thr Ser Arg
20 25 30

Pro Gly Thr Gly Leu Arg Cys Arg Ser Cys
35 40

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Pro Arg Ala Tyr His Met Val Cys Leu Asp Pro Asp Met Glu Lys Ala
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Pro Glu Gly Lys Trp Ser Cys Pro His Cys
35 40

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<400> 6

Cys Arg Val Cys Lys Asp Gly Gly Glu Leu Ile Cys Cys Asp Thr Cys
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Pro Ser Ser Tyr His Ile His Cys Leu Asn Pro Pro Leu Pro Glu Ile
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Pro Asn Gly Glu Trp Leu Cys Pro Arg Cys
35 40

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1 5 10 15

Pro Lys Val Phe His Leu Ser Cys His Val Pro Thr Leu Thr Asn Phe
20 25 30

Pro Ser Gly Glu Trp Ile Cys Thr Phe Cys
35 40